SEQUENCE LISTING

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::110 . Poche Vitamins AG
    -120 Microbial process for producing L-ascorbic acid and
           :-erythorbic acid
    -130 - Alicyclobacillus NA20, 21, FJ21 16S nuc
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    +150 - EP Application No. 00118059.5
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    ×160× 3
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    -311/ 1529
    <212 - ENA</pre>
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D.
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    decoyuaataa egodeggaaa egggegetaa tgeoggatae geoegegagg aggeatette 130
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13.
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    quanticiant integrated analogy tag against graph 	ag{2.0}
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50
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55
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   · 211: 1529
    +2120 DNA
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    toopquatoa otgogoqtaa aggytgogta ggoggtogag caaytotoga gtgaaagtoo 🕬
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    quaecttacc agggettgac atocstetga caccetsaga gatgaggggt costtogggg 1920
   cagaggagac aggtggtgca tggttgtcgt cagstcytgt cgtgagatyt tgggttcagt 1930
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11
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4,31
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35
    <210+3
<211+1495
<210+DDA
    #212 Alievelobacillus sp.
    +1111 + SENA
    %1.12 * 1)..(1495)
%7.13 * FU-21 Partial 16SrFNA gene sequence
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    aangagagit aatgeoggat aegeoogega ggaggeatot tettgeggeg aaaggeooga 1\%
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    aaggregacya tgogtagoog acctgagagg gtgacoggod ababtgggab tgagacabgg 3\%
    coda jastoo taogggaggo agoagtaggg aatottoogo aatgggogoa agootgaegg 383
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    eggrateggg agtggalage edeatgegag aeggtabega gtgliggaage eeeggcthae
    tamm's group dagoogogyt aaaaogtagg gggogayogt tgtooggaat bactgggogt
    aautygygyg taggogytog agbaagtotg gagtgaaagt boatggotba abbatgggat \delta^{\circ} .
    gg. Ttqgaa actgottgae ttgagtgotg gagaggoaag gggaattooa egtgtagogg 6\%
    tgalastgojt agagatgtgg aggaatabda gtggdgaagg ogodttgotg gabagtgabt
    gathithaly cacgaanged tggggagbaa acaggattag atacotiggt agtocacgeb 78
   gt:swogaty agtgotaggt gttgggggga cacaccecag tgccgaagga aacccaataa 84\%
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    casaagtagt ggagcatgtg gtttaattog aagcaacgog aagaacotta ccagggottg 90
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